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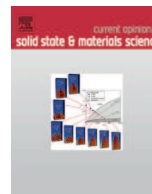
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High throughput screening for discovery of materials that control stem cell fate

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ABSTRACT

Insights into the complex stem cell niche have identified the cell–material interface to be a potent regulator of stem cell fate via material properties such as chemistry, topography and stiffness. In light of this, materials scientists have the opportunity to develop bioactive materials for stem cell culture that elicit specific cellular responses. To accelerate materials discovery, high throughput screening platforms have been designed which can rapidly evaluate combinatorial material libraries in two and three-dimensional environments. In this review, we present screening platforms for the discovery of material properties that influence stem cell behavior.

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1. Introduction

The ability of stem cells to self-renew or to differentiate into specialized progeny makes them a valuable source for production of clinically relevant cells for regenerative medicine, disease modeling and biomedical applications. Stem cells broadly fall into two categories. The first, human pluripotent stem cells (hPSCs), include embryonic stem cells (hESCs) and have the potential to generate cells from any of the three germ layers that comprise all of the ~200 cell types found in the body [1]. Also included in this group are induced pluripotent stem cells (hiPSCs), which bypass the need for cultivation from a blastocyst by reprogramming somatic cells into a stem cell state using a cocktail of transcription factors [2]. The second group encompasses tissue specific or ‘adult’ stem cells whose role is to assist in repair or renewal of tissue. These cells are generally considered multipotent meaning that their differentia-

tion potential is limited to the cell types of the tissue in which they reside.

The promise of stem cells in regenerative medicine is becoming reality with recent approval for the use of limbal stem cells for the treatment of ocular burns [3] and phase I clinical trials underway for the use of hPSC derivatives for spinal cord injury [4] macular degeneration [5] and heart failure [6]. To broaden the application of stem cells and their derivatives for wide ranging conditions there is a need for culture systems that enable controlled manipulation of these cells.

The first successful *in vitro* propagation of hESC was accomplished in 1998, this was over a decade after the culture of mouse embryonic stem cells (mESC) was achieved [7]. The culture conditions found to maintain mESC pluripotency could not be translated to the human counterparts, where pluripotency could only be maintained when the cells were cultured on a feeder layer of mouse embryonic fibroblasts (MEFs) [1]. It was later discovered that this MEF layer could be replaced with a basement membrane matrix extracted from mouse sarcoma cells such as Matrigel™ [8]. While these advances have enabled the culture of pluripotent stem cells outside of the body, the field requires culture conditions that are primed for clinical translation such as those that are scalable,

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defined, reproducible and xeno-free. High throughput screening strategies have been adopted to search for substrates that achieve these goals [9].

The critical role of the supporting substrate in maintaining pluripotency of human stem cells *in vitro* has been apparent since their derivation. Growth substrates to recapitulate the extra cellular matrix (ECM) such as Matrigel™ or its components, such as laminin, have been commonly used [8]. More recently, substrates bearing epitopes that are capable of interacting with cells have been developed, for example Synthemax™ is an acrylate substrate conjugated to RGD peptide derived from vitronectin that can support self-renewal of hESCs [10]. The RGD ligand is a cell adhesive peptide that interacts with cell surface integrins [11]. Integrins and other cell adhesion molecules (CAMs), such as cadherins have all been implicated in regulating cellular behavior from maintaining pluripotency to directing differentiation [12]. Advances in the characterization of stem cell interaction with their environment has demonstrated that material physicochemical properties including chemistry, topography, geometry and stiffness also play an active role in modulating stem cell fate, particularly demonstrated with mesenchymal stem cells (MSCs) [13–16] (Fig. 1).

In the body, stem cells reside in a complex niche and receive a multitude of cues from the surrounding ECM, cell–cell contact and soluble factors contained within the aqueous milieu. In addition, the same stimuli may trigger a different biological response depending on the stem cell type. This and other complex structure–function interrelationships, some of which are not fully known, hinder a rational approach in the design of stem cell culture substrates, as it is difficult to predict how a given material property or combinations thereof will bias stem cell fate. More recently, the discovery of naïve states of hESCs and the difficulty in optimizing their culture conditions emphasizes the need for methods to keep pace with the rapidly evolving field [17]. Therefore, researchers have adapted high throughput screening (HTS) strategies to identify culture substrates that are appropriate for stem cell culture [9]. HTS has been utilized in a pharmaceutical setting facilitating early stage drug discovery since the 1980s. Libraries of compounds can be assayed for activity against a biological target to generate lead candidates essentially when structure based design is not possible. Such approaches rely on innovation in robotics for automation, robust biological assays to minimize false positives and high content analytical tools [18]. Adoption of the HTS strategy to accelerate the discovery of materials that can direct stem cell fate began around a decade ago [19,20]. By applying combinatorial methods used in conventional HTS, the

structural diversity of polymer libraries can be exponentially increased [20]. In addition, the design of material libraries can be guided by the outcome of biological activity. For this, a suite of high throughput materials characterization techniques is also required to generate comprehensive datasets that can be correlated to biological activity using statistical methods that identify structure activity relationships (SARs) in a systematic and unbiased fashion to enable a more rational approach to optimize materials identified from such screens [21–23] (Fig. 2).

Synthetic materials allow for greater manipulation and control of physical and chemical properties compared to biological substrates, lending to design of modular systems that can be simplified to uncouple substrate effects. In addition, for clinical applications, consistent material quality and function can be assured with fully characterized synthetic substrates, however it remains to be seen if these materials can recapitulate the complex nature of biological matrices. Nonetheless, HTS strategies can help to discover influential material properties to feedback into the design of robust differentiation systems, aid the isolation of rare or difficult to culture cell populations and begin to unravel complex molecular pathways underpinning the identified cell–material interaction.

This review will focus on an overview of the HTS systems designed to probe the interaction between material properties and stem cell phenotype, including surface chemistry, topography, elasticity and 3D micro-environments.

2. Substrate chemistry

Tissue culture polystyrene (TCPS) is a widely used synthetic growth substrate suitable for various cell types including human mesenchymal stem cells (hMSCs). However, simple substrates such as TCPS have limited cellular interaction and usually require coating with ECM proteins and/or soluble factors from the culture medium to modulate the behavior of adherent cells [24]. This has led to the development of a new wave of synthetic growth substrates that have a broad range of surface chemistries to elicit a particular cell response. The surface chemistry of materials has been used to achieve the desired biomolecular adsorption from the culture medium to control cell response and/or act in itself as a ligand for cellular interaction. HTS of proteins, peptide fragments or chemical moieties presented at the substrate surface, to invoke a desired response (e.g. maintaining pluripotency or directing differentiation toward a specific lineage) have been widely explored and will be discussed in this section.

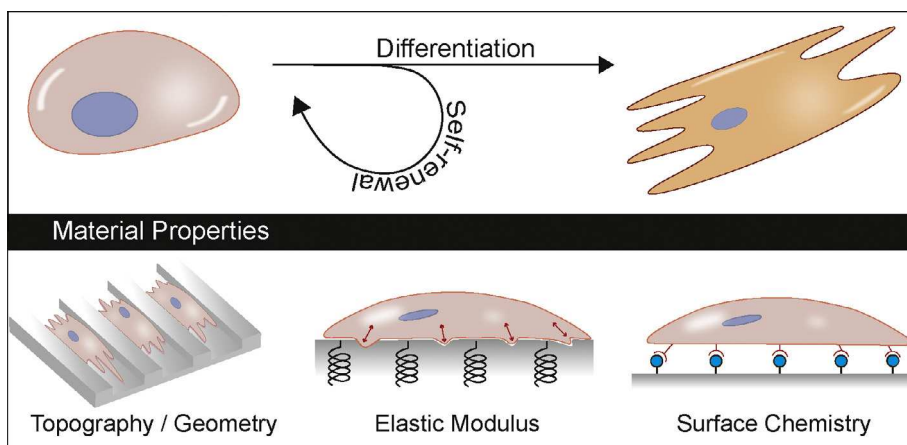


Fig. 1. The culture substrate that stem cells adhere to can harness material properties such as topography, patterning, elastic modulus, surface chemistry and combinations thereof, to influence stem cell fate.

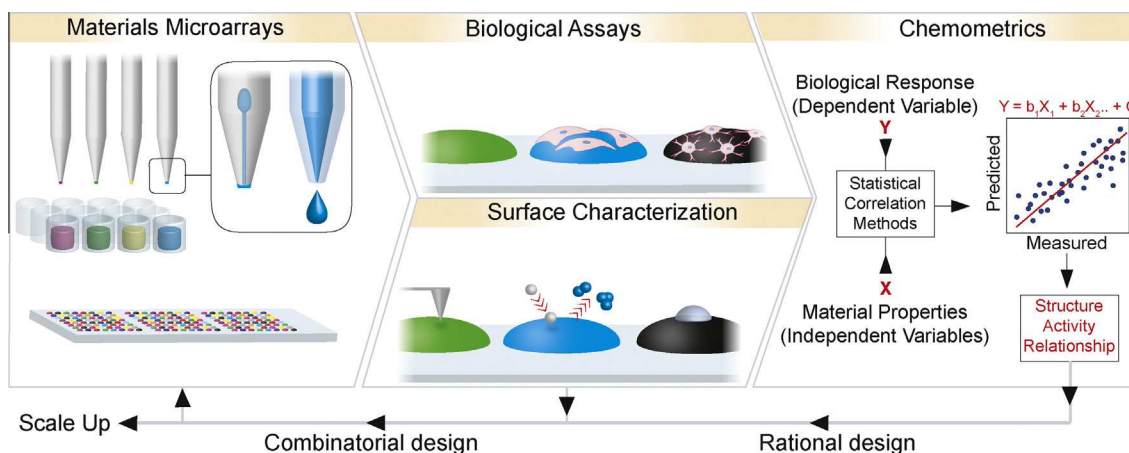


Fig. 2. Materials microarrays can be fabricated by automated robotics including contact and ink-jet printing shown inset in first panel, left and right, respectively. Characterization of cell response and corresponding surface properties generate large data sets that can be correlated to define structure activity relationships that inform the rational design of biomaterials. For this purpose, all data is input into statistical modeling to identify positive and negative relationships. However, for 'hit' material generation, the best performing materials can be taken forward to fabricate combinatorial libraries of materials to be investigated in further iterations of HTS platforms.

2.1. Proteins and peptides

HTS studies targeted toward understanding the necessary biomolecules that direct stem cell fate were focused on ECM proteins. Solutions of fibronectin, laminin and collagen I, II and III in various combinations were robotically spotted onto acrylamide-coated slides [25]. Adhered mESCs cultured on these arrays were confined to the ECM environments and assayed for expression of an early hepatic fate marker, β -galactosidase, to identify conditions that encourage hepatic differentiation. Several hepatic promoting conditions were identified with nine of the ten highest signals measured on cells adhered to collagen I containing matrices.

Smaller epitopes, such as peptides, can be easily handled and modified for immobilization using common chemistry techniques to increase the throughput of these screens. Surface modification strategies such as self-assembly are powerful tools to immobilize ligands capable of binding to cell surface integrins. The high spatial resolution with which self-assembling monolayers (SAMs) can be generated enables discrete chemical moieties to be screened in parallel (Fig. 3A). To facilitate SAM array preparation, gold surfaces have been modified with perfluoroalkane thiols that can be subsequently removed via UV irradiation through a photo-mask to produce high-resolution patterns. Arrays of peptide-substituted alkanethiols can then be prepared by spotting multiple solutions within the generated pattern, the fluorinated SAM prevents spreading of the spotted peptide-thiol solutions and also creates a low bio-fouling surface for cell screening. Phage display was employed for initial identification of cell binding peptides that could then be presented as a SAM to identify which binding peptides could also be used as a cell supportive substrate. Around 30,000 peptide-presenting phages were identified that were able to bind embryonic carcinoma (EC) cells. This initial pool was reduced to seven peptides with significant cell binding potential over peptide-free phages using a cell suspension enzyme-linked immunosorbent assay (CS-ELISA). When immobilized as a SAM two of these peptides, TVKHRPDALHPQ and LTTAPKLPKVTR, were able to support human embryonic stem cell (H9 cell line) growth at similar levels to Matrigel™ for 20 days (3 passages) [26]. Interestingly, cell adhesion was not significantly reduced by addition of EDTA or heparin suggesting that adhesion to the phage-derived peptide surfaces was not mediated by integrins or proteoglycans. This method of screening peptide motifs demonstrates the utility of HTS to identify novel functional peptides that could not have been predicted from what is currently known.

A similar approach has been applied to screening peptides that contain RGD and glycosaminoglycan binding epitopes, the most successful of which being a heparin-binding peptide derived from vitronectin (Table 1) [27]. At densities of 0.5–25%, this peptide was able to support self-renewal of hiPSCs and hESCs when combined with Rho-associated kinase (ROCK) inhibitor or cyclic RGD peptide. More recently, surfaces bearing this peptide in combination with RGD containing peptides have been used to promote ectoderm and neuronal differentiation. The cues for cell differentiation from this substrate were attributed to cell surface integrin engagement and subsequent stimulation of the Akt signaling pathway via integrin linked kinase [28]. Biotinylated-peptides were immobilized to streptavidin-coated TCPS dishes but scalability over large areas was not demonstrated limiting the applicability of using such peptides for stem cell expansion. SAMs have also proved to be a useful tool in investigating the relationship between surface chemistry mediated-adhesion protein binding and MSC differentiation [29].

To ease the manufacture of generating combinatorial peptide arrays, 384-well plates were prepared containing mixed solutions of alkanethiols bearing azide functionality and peptides modified with alkyne functionality. Conjugation of the peptides to the alkanethiols was achieved using azide-alkyne "click" chemistry prior to spotting the solutions onto gold surfaces. Combinatorial mixtures of cell adhesion peptide, a bone morphogenetic protein 7 (BMP7) growth factor derived peptide, and a heparin binding peptide were printed [30]. The adhesion peptides allowed long-term (>1 week) observation of cell behavior with BMP peptide directing osteogenic differentiation of adipose derived stem cells. This was determined by the increased expression levels of osteogenic markers runt-related transcription factor 2 (Runx2) and osteopontin (OPN) from cells cultured on BMP peptide surfaces versus adhesion peptide only controls.

Recently, a combinatorial bimolecular nano-patterned platform was designed to study the effects of cell behavior on nanoscale topography, controlled biomolecule conformation and identity in parallel over large areas (mm^2 per pattern). Vitronectin and Laminin 1 were mixed in 8 different ratios and immobilized on 8 nanoscale dimensions (100 nm – 150 μm) in duplicate to create 128 combinatorial dual protein patterns on a single surface. Adhesion profiles of human dental pulp stem cells were determined across the environments. Line widths less than 500 nm encouraged focal adhesions and spreading across the patterns. However, widths greater than 700 nm guided adhesion and spreading along the patterns identifying the importance of nanoscale geometry on cell adhesion which will be discussed in further detail in Section 3 [31].

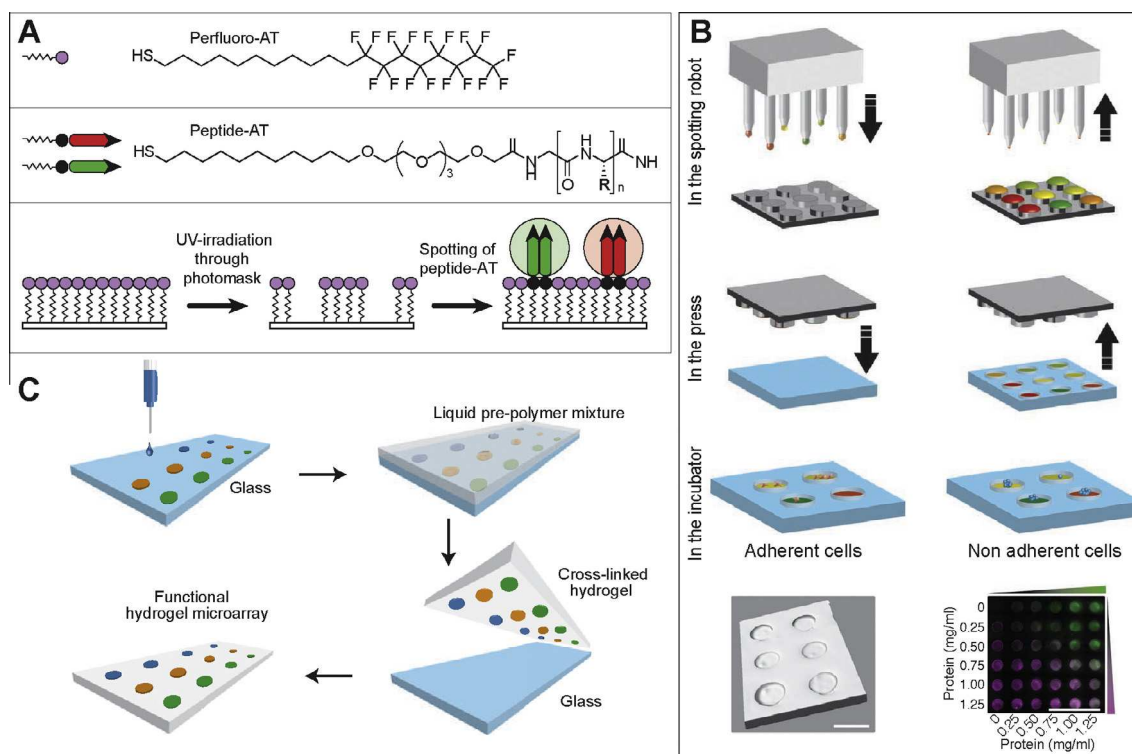


Fig. 3. High throughput surface modification strategies to probe stem cell behavior. (A) Self assembling monolayers of perfluoroalkane thiols can be photo-patterned to allow spotting of peptide-substituted alkanethiols (AT) presenting ligands for cellular interaction. (B) Microwells of unique protein microenvironments can be fabricated using contact printing to deposit solutions onto a silicon stamp to subsequently press into a PEG substrate to fabricate artificial niches. (C) Hydrogel microarrays are fabricated by first depositing substrates onto glass and which is embedded under a thiolated PEG and bis-acrylated PEG mixture that is UV cross-linked in the presence of a photoinitiator. The PEG hydrogel is peeled from the glass layer to expose the functionalized PEG surface. Adapted with permissions, (A) [26] 2010© ACS, (B) [46] 2011© NPG, (C) [45] 2009© NPG.

Table 1

Peptide sequences found via HTS platforms to maintain pluripotency or promote differentiation.

Peptide sequence	Origin	Function	References
TVKHRPDLHPQ	Phage display	Maintain pluripotency	[26]
LTTAPKLPKVTR	Phage display	Maintain pluripotency	[26]
GKKQRFRHRNRKG	Vitronectin	Maintain pluripotency	[27]
YIGSR	Laminin	Cell adhesion	[30]
KPSSAPTQLN	Bone morphogenetic protein 7	Osteogenic differentiation	[30]
KRSR	Heparin-binding protein	Cell adhesion	[30]

2.2. Polymers

Synthetic polymers offer an alternative to biological epitopes for the control of stem cell fate. The use of polymer microarrays to screen for their influence on the propagation of hESCs was achieved by Anderson et al. in 2004 and by applying principles of combinatorial materials design [32,33], demonstrated rapid synthesis of 576 acrylate co-polymers from a library of 24 monomers mixed in binary 70/30 % v/v ratios. Polymer microarrays are a powerful tool on which thousands of materials can be investigated in parallel on a single slide for desired cell responses [19,20,34]. Array fabrication can be performed using automated robotics. Ink-jet printing offers more flexibility and automation to enable easy drop volume control and drop in drop mixing of reagents to bypass manual pre-mixing. However, the wetting behavior and viscosity of the monomer solution highly influence droplet formation

therefore, uniform deposition of solutions with wide ranging surface energies or viscosities can be challenging. Contact printing generally affords greater control over spot size because it is predominantly the geometry of the pin that determines the size of polymer spot enabling a wider range of solution viscosities to be printed uniformly [21] (Fig. 2). Once optimized, both techniques enable rapid array fabrication of large numbers of polymers printed in discrete and identifiable regions. A strength of the microarray platform relative to casting polymers into microwells is the reduction in scale of materials required which transform impracticably large, laborious and expensive experiments into efficient and economical assays. To prevent background cell and bio-molecule adhesion between polymer spots that may lead to cross-talk, a bio-resistant polymer is used to coat the substrate that is to be printed upon such as poly(2-hydroxyethyl methacrylate) (pHEMA) [19], acrylamide [35] or poly(ethylene glycol) [36]. However, due to this lack of spatial separation, it should be noted that any paracrine factors secreted by cells are free to diffuse through the shared culture medium for interaction with cells adhered to neighboring polymer islands. To partially abate such effects, polymer islands can be printed in replicate and in randomized regions on the array.

Several groups have employed polymer microarrays as a means to discover suitable substrates to address a specific need. For purposes of gene targeting in stem cells, low seeding densities are required to allow clonal growth from single cells that have been correctly targeted. Mei et al. screened polymer microarrays for substrates that could support clonal propagation of hESCs from single cells [37]. Arrays were fabricated from 16 major monomers and 6 minor monomers mixed at 6 different ratios to generate 496 unique materials. A significant advance in this study was to perform the screen using one of the first commercially available

media, mTeSR1, made up of relatively defined components compared to conventional MEF conditioned medium. However, arrays required coating with fetal bovine serum (FBS) to overcome poor cell attachment. In an attempt to further refine the culture system, the authors investigated replacement of the FBS coating with three proteins, albumin, laminin or vitronectin. Integrin blocking experiments revealed that hit polymers enabled $\alpha_v\beta_3$ and $\alpha_v\beta_5$ integrin binding of hESCs to adsorbed vitronectin and although vitronectin coated TCPS is known to support hESC growth, not all investigated polymer substrates coated with vitronectin supported clonal growth. This work was extended to quantify how roughness, elastic modulus, wettability and surface chemistry of the microenvironment regulate self-renewal in ESCs concluding that polymer surface chemistry plays a dominant role.

Microarrays have also facilitated discovery of a poly(4-vinyl) phenol substrate that can support proliferation of hPSC derived neural progenitor cells [38] and a methacrylamide polymer that can maintain hPSC pluripotency for over 20 passages in mTeSR1™ medium. Using quartz crystal microbalance (QCM) experiments it was found that bovine serum albumin (BSA) was critical for cell adhesion to the polymer [39]. However, BSA-containing media such as mTeSR1™ and StemPro® although better defined than MEF conditioned medium, may still contain other undefined factors that bind to BSA (e.g., Immunoglobulin E (IgE) antibodies), which can confound results in assays that may vary from batch to batch [40]. Developments in the propagation of hPSCs in fully defined media such as serum-free and xeno-free E8 medium [41], which contains only 8 components including transforming growth factor beta (TGF β) and fibroblast growth factor 2 (FGF2) aids unraveling the precise role of isolated material properties without confounding sources of unsolicited cues from, for example FBS and BSA. The prospect that simplified culture media such as E8 can be used reproducibly to maintain pluripotency and generate terminally differentiated cells provides encouragement to the materials science community that physicochemical cues from substrate materials can be developed to differentiate stem cells “on-demand” through stimuli supplied via substrate chemistry [42].

To demonstrate the ability of materials found by HTS to not only propagate hPSCs but to also support media induced differentiation of all three germ layers, Celiz et al. investigated the largest starting library to date of 141 (meth)acrylate and (meth)acrylamide monomers in multi generational contact printed arrays of evolving libraries based on previous hit materials. Three generations of arrays were investigated, the first to identify homopolymers that could support hPSC in Stempro medium, the second co-polymerized hits in major/minor ratios and the third expanded different ratios of hit combinations identified from the second generation to optimize cell response. Using this approach, 909 different substrates in 4356 individual assays were analyzed and lead to the discovery of a co-polymer poly(HPhMA-co-HEMA) that could be scaled up to coat common culture-ware, support hPSC expansion through 5 serial passages and allow directed differentiation to cardiomyocytes, hepatocytes and neural cells [43].

With the scale of HTS increasing, automated processes are required to assist fabrication of these platforms. Hansen et al. recently demonstrated the capability of ink jet printing for *in situ* mixing to facilitate rapid combinatorial polymer preparation [44]. Varying ratios of co-polymers were generated from a library of 26 acrylates and acrylamides. Elegant drop in drop mixing of monomers, photoinitiator and cross-linker was achieved to fabricate microarrays comprising of 7316 polymer islands. The low-fouling substrate was synthesized using a two-step fabrication procedure consisting of arrayed aqueous sucrose solution to serve as a mask followed by immersion in a cytophobic fluorinated silane. Removal of the mask and subsequent immersion in a methacrylated silane enabled the arrayed biomaterials to resist

spreading prior to photopolymerization. This screening platform was able to identify a polymer, a copolymer of 2-(dimethylamino)ethyl methacrylate and ethylene glycol dicyclopentenyl ether acrylate, that could maintain pluripotency of RH1 cells (hESC) for 5 passages in mTeSR1™ medium.

The fabrication of microarrays on softer substrates with a Young's modulus in the kPa range, that is more physiologically relevant than the GPa moduli displayed by conventional glass slides, has been demonstrated by Gupta and co-workers [45]. This was achieved by arraying thiol- or alkene-functionalized biomolecules onto a glass substrate that was subsequently coated with a liquid PEG pre-polymer. Curing via UV irradiation enabling coupling of the arrayed biomolecules to the PEG hydrogels which were peeled off the glass substrates to expose the hydrogel microarray (Fig. 3C). The orthogonal nature of thiol-ene chemistry was used to immobilize a large selection of molecules at the surface of the PEG hydrogel.

Thus far, while chemical moieties have been identified that can improve maturity of hESC derived cardiomyocytes in serum-free medium [47] and sustain the differentiation of hPSC derived hepatic progenitors to hepatocytes using a maturation medium, no polymer has been shown to initiate the formation of the three germ layers, or to completely replace a biological substrate in the multi-step differentiation process from pluripotent to terminally-differentiated cell [48]. The identification, characterization and scale-up of such a substrate are major challenges for future investigation.

3. Material physical properties

In addition to the biochemical cues discussed earlier, stem cells also respond to biophysical cues from their surrounding environment. Forces exerted via cues such as stiffness, topography, ligand density and stretch lead to activation of intra-cellular pathways via focal adhesions that connect the external ECM to the cellular cytoskeleton. This conversion of biophysical cues to biochemical signals is termed mechanotransduction [49]. Significant progress has been made in uncovering some of the transducing pathways relevant to guiding stem cell fate decisions. For example, phosphorylation of focal adhesion kinase, has been implicated in modulating response of hMSCs to nanotopography [50]. Whereas, the RhoA-ROCK signaling pathway was shown to be activated by geometric cues that influence hMSC shape and spread which subsequently determined adipogenic (rounded, less spread shape) or osteogenic (spread) commitment [15]. However, without prior knowledge of precisely defined ECM-cellular interaction it is difficult to predict how discrete presentations of biophysical cues determine cell fate. This section will focus on physical material properties that are amenable to HTS platforms such as surface topography and substrate stiffness to aid investigation of biophysics on cellular behavior and guide the design of biophysical features.

3.1. Topography

Topographically patterned substrates have been shown to direct stem cell fate of MSCs [51], hESCs [52,53] and hiPSCs [54]. Dalby et al. used electron beam lithography to precisely fabricate nanopits of 120 nm diameter and 100 nm depth arranged in 5 different patterns to compare varying degrees of disorder with the aim to re-create the natural disorder observed in the ECM protein, collagen. They fabricated polymethylmethacrylate (PMMA) arrays embossed with the various patterned topographies and found that a disordered topography, where nanopits 300 nm apart in a square array were randomly displaced by up to 50 nm, was able to induce

osteogenic differentiation of hMSCs that was comparable to cells that were chemically induced using dexamethasone on planar substrates. Interestingly, using ingenuity pathway analysis and canonical analysis the authors reported signaling events that were specifically triggered when cells were cultured on the patterned surfaces including fibroblast growth factor signaling and epithelial growth factor signaling which are associated with bone development [51]. Applying such in depth analysis to HTS provides a means of uncovering complex pathways of successfully identified material properties. The effect of topographical disorder on hESCs was later investigated using the same nanopit topography hot embossed onto polycarbonate. The study revealed that mesodermal lineage specification of hESCs could be encouraged using topographical cues alone in basal medium. Despite the observation that planar surfaces could also direct mesodermal differentiation without chemical factors, the authors report that the additional topography enhanced up-regulation of a panel of mesenchymal specific markers that could give rise to mesodermal derivatives such as bone and cartilage and down regulation of non-mesenchymal markers [53].

Ankam et al. investigated the influence of topography on glial or neuronal specification during neural differentiation of hESC. They fabricated a multi-architecture (MARC) chip using nano-imprint lithography that was used as a master mold to pattern PDMS. The 2×2 cm MARC chip is assembled with 18 topographies in duplicate (Fig. 4A). On planar surfaces, hESC differentiation in a neuro-basal medium (N2B27) resulted in a mixture of neuronal and glial cells. A higher neuronal population was found on anisotropic lines whereas culture on isotropic patterns resulted in a greater proportion of the glial sub-type [52].

For exploration of a large topographical design space, the de Boer group developed a mathematical algorithm that could design over 150 million different topographies by combinatorial mixing of

three primitive shapes. Shapes ranging from 3 to 10 μm were arranged to form features ranging from 10 to 28 μm and 5 μm in height (Fig. 4B.i). From this library 2176 features were selected to make a silicon master etched using photolithography. Poly(lactic acid) (PLA) films were patterned with the master using hot embossing to fabricate 'Topochips'. Cell material interactions were screened using high content imaging to identify topographical patterns that were able to maintain pluripotency of hMSCs or induce osteogenesis (Fig. 4B.ii) [55]. Identification of such topographies will enable future investigation into the mechanisms underpinning their ability to control stem cell fate.

3.2. Substrate stiffness

Cells are continuously assessing substrate stiffness through mechano-sensing and similar to soluble factors, matrix stiffness can influence stem cell lineage specification [56]. Engler et al. elegantly highlighted the profound effect that substrate stiffness can have on stem cell fate by culturing MSCs on polyacrylamide gels of three physiologically relevant moduli that mimicked brain (1 kPa), muscle (10 kPa) and bone (100 kPa) for 1 week. It was found that cells displayed morphology and RNA profiles indicative of neurogenesis, myogenesis and osteogenesis respectively [57].

To address the need for a high throughput platform that would allow multifactorial investigation of varying substrate stiffness and biochemical composition on stem cell differentiation, Gobaa et al. fabricated soft hydrogel microwell arrays where the protein micro-environment of each well could be varied [46]. Protein was deposited by contact printing onto each micro pillar of a silicon stamp and then pressed into a thin film of PEG hydrogel to create individual microwells of a confined protein niche. The proteins were tethered using thiol terminated PEG macromers in the hydrogel which were available to react with maleimide functionalized

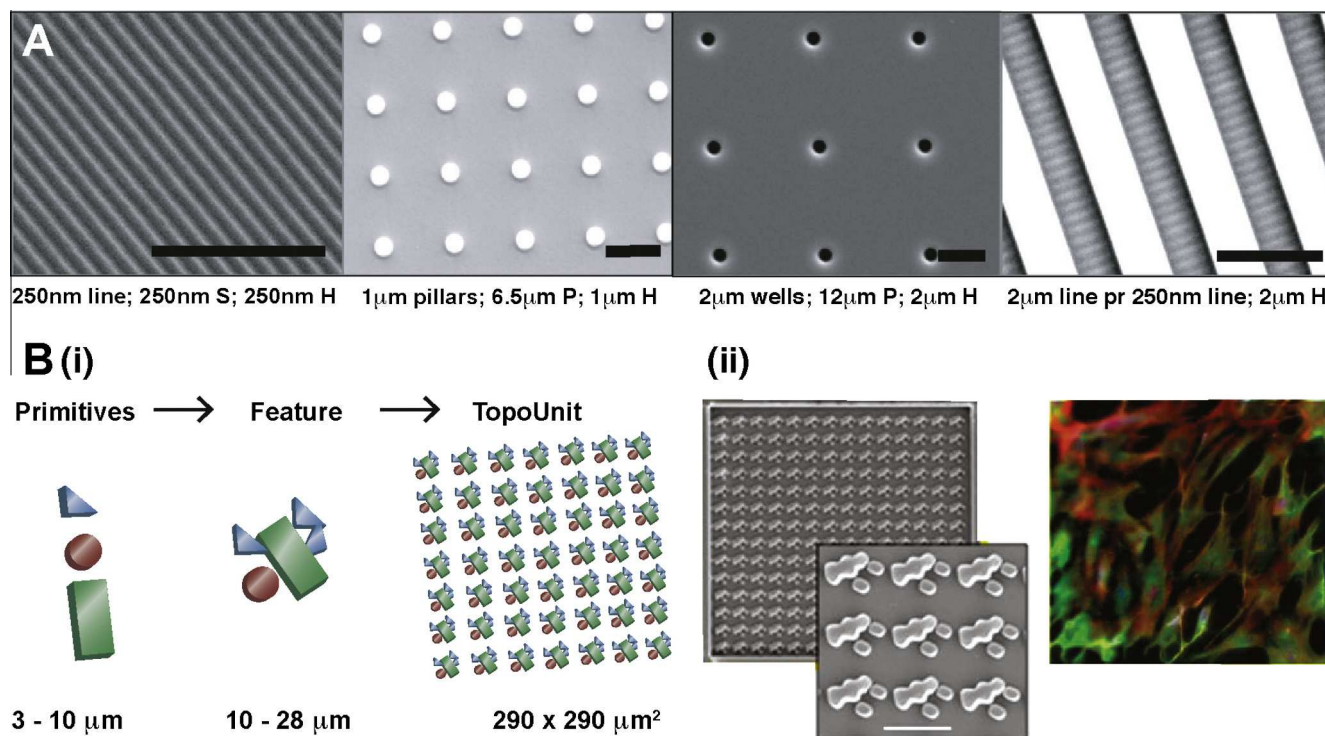


Fig. 4. Topographical assays developed for stem cell screening. (A) Scanning electron microscopy images (SEM) show patterned PDMS. The geometries were transferred from master molds using soft lithography and used for investigation of hESC neural differentiation. (S = spacing, H = height, pr = perpendicular). (B) i. TopoChip pattern generation using combinatorial design with three types of primitive shapes. ii. SEM of one TopoUnit pattern. hMSCs were cultured on this topography and displayed the highest immunostaining for alkaline phosphatase, a marker for early osteogenic differentiation. Scale bar = 20 μm , height of feature is 5 μm . Reproduced with permissions, (A) [52] 2013©, (B) [55] 2011©.

proteins. Using this strategy an array with dimensions of a standard slide, 2.5 cm × 7.5 cm, could contain up to 2016 microwells that could be seeded with cells in a single well of a rectangular four well plate (Fig. 3B). The authors found that the microarrays could be reproduced on PEG hydrogels with a Young's modulus ranging from 3 to 150 kPa. Stiffness was achieved by adjusting the ratio of thiol and vinyl terminated PEG macromers, due to this the authors note that gels of varying stiffness also changes the number of thiol groups available for the maleimide protein conjugation which will vary the density of ligand presentation and that the molar concentration of free groups for ligand functionalization should be kept constant to overcome this. The study used the hydrogel microwell array to investigate the influence of substrate stiffness on hMSC differentiation and found that increasing the stiffness of the substrate increased osteogenic differentiation.

Mih et al. fabricated polyacrylamide arrays of varying substrate stiffness's that were spatially separated by casting the gels in glass bottomed 96 and 384 multi well plates. A range of Young's moduli from 0.3 to 55 kPa could be investigated in one multi well format by varying the bisacrylamide cross-linking ratio. The assay format was used to demonstrate applications for analysis of 7 different cell lines including hMSCs [58]. The authors aimed for PA films that were at least 3 times the threshold 20 μm thickness that has been reported to prevent MSCs from sensing a rigid polystyrene substrate underlying a 1 kPa PA coating [59].

4. Materials for surveying 3D micro-environments

While 2D platforms have been indispensable in constructing an understanding of stem cell fate control outside of the body, the native stem cell niche is inherently three-dimensional (3D). Recently, it has become clear that cell function often depends directly on the dimensionality of their surrounding matrix and that studying biology exclusively in 2D is insufficient. For example, freshly isolated murine hepatocytes de-differentiate when plated on tissue-culture polystyrene (TCPS) and fail to maintain liver-specific function [60]. In addition, while it is known that hMSC differentiate toward an osteogenic lineage on stiff 2D substrates [61], it has been observed that stiff and non-degradable matrices favor an adipogenic lineage in 3D [62]. In this work, adipogenesis was favored because cells were not able to spread; whereas, osteogenic differentiation was favored in degradable 3D matrices as they enabled hMSCs to spread and generate contractile force, which was necessary for osteogenesis and echoes the response to biophysical geometric cues in 2D reported by McBeath [15]. Yet, all of the attributes of the micro-environment that discriminate 3D and 2D culture from the viewpoint of the cell still remain unknown.

2D environments present a highly polarizing environment with monolayer ligand presentation in which only a fraction of the cell membrane can interact with the surrounding ECM and other cells while the remainder of the cell is exposed to bulk culture media [63]. This diverges from the native ECM in which cells are immersed in a fibrous network of proteins with sequestered and diffusing growth factors as well as interactions with other cells in all directions via cell–cell contacts or paracrine signaling [64]. Individual signals within the complex signaling milieu of the ECM combine synergistically and antagonistically in a non-linear manner [65]. Therefore, 3D HTS techniques enable researchers to understand more completely the bidirectional and multimodal interactions between cells and the 3D ECM in a systematic manner.

Many 3D cell culture platforms are composed of natural protein hydrogels (e.g., collagen or Matrigel™) into which cells can be encapsulated. These fibrillar protein matrices are useful but they are challenged by batch-to-batch variability while providing the

user little control over individual cues such as ligand presentation, elasticity, and topography. Fortunately, the toolbox for 3D cell culture has expanded significantly in the recent years as researchers have developed synthetic hydrogels with tunable mechanical and biochemical properties [66]. Much of this work has focused on the covalent cross-linking of hydrophilic polymers in the presence of cells. Polymer and cross-linking density control hydrogel mechanics while biochemical ligands can be incorporated into the network as whole proteins or peptide mimics [67]. Encapsulation of cells within synthetic hydrogels is often achieved using photopolymerization via chain polymerization of (meth)acrylates or thiol–ene click chemistry [68,69]. Highly efficient, water soluble, and cytocompatible photoinitiators exist for rapid fabrication of gels with visible light [70]. In parallel, several bio-orthogonal chemistries (e.g., copper-free click chemistry or tetrazine–norbornene coupling) have been utilized for the fabrication of 3D ECM mimics that proceed without a catalyst at ambient conditions in cell culture medium [71,72]. In each case, a liquid precursor of polymeric substituents, biochemical ligands, and cells is transformed into a gel for the 3D encapsulation of cells.

While rationally-designed 3D biomaterials have been used predominantly in systematic studies of individual cell–matrix interactions they are generally amenable to HTS. Sala et al. leveraged liquid handling robotics to form arrays of artificial extracellular matrix (αECM) through the selective, enzyme-mediated crosslinking of peptide-functionalized PEG to screen cell function within a range of αECMs [73]. The complexity of cell–matrix signaling requires experimental design that investigates non-linear interactions between multiple cues, which even with a limited number of inputs, generates a large number of experimental conditions, making HTS an attractive strategy.

Robotic handling of living cells presents unique challenges. Critically, the precursor solution (viscosity, chemical composition, and additives), cross-linking strategies, as well as the handling techniques must not be cytotoxic or mutagenic. Pioneering groups have developed a range of suitable 'bio-inks' (biologically-compatible printing solutions) based on advances in the field of 3D stem cell culture using auto-gelation, photoinitiation, or thermal gelation for the fabrication of 3D microarrays. Notably, Lutolf and coworkers pioneered the fabrication of modular 3D hydrogel arrays using liquid handling robotics to investigate the dynamic interplay between mechanical and biochemical signals in 3D ECM mimics [74] (Fig. 5A). Specifically, the authors created defined microenvironments through selective enzyme-mediated crosslinking of factor XIIIa (FXIIIa) with PEG modified with substrates for FXIIIa. Additional modification of PEG to include protease susceptible peptides, enables degradation and remodeling similar to native environments through matrix-metalloproteinase (MMP) cell secretion. This approach enables defined control over initial mechanical properties (E, Young's moduli) by tuning polymer concentration; degradability by altering the MMP-sensitive peptide linkage; cell density down to single cells; ECM components (e.g., collagen IV, fibronectin, laminin); cell–cell signals (e.g., E-cadherin, Jagged, EpCAM); and soluble factors (e.g., FGF4, BMP4, LIF). Additionally, this strategy facilitates real-time monitoring of cell function as the microgels are optically clear and can be formed in standard 1536-well plates for complete isolation of the 1024 possible unique microenvironments. Oct4-GFP expressing mESCs were encapsulated within each microenvironment for facile readout of mESC pluripotency. Systematic analysis of the effects of microenvironment on mESC proliferation and self-renewal was achieved using generalized linear models (GLM). The initial screen demonstrated that soluble factors (FGF4, BMP4, and LIF) were the strongest predictors of mESC self-renewal and proliferation with an additional strong effect of matrix mechanics. Gels with low elasticity led to increased self-renewal and proliferation.

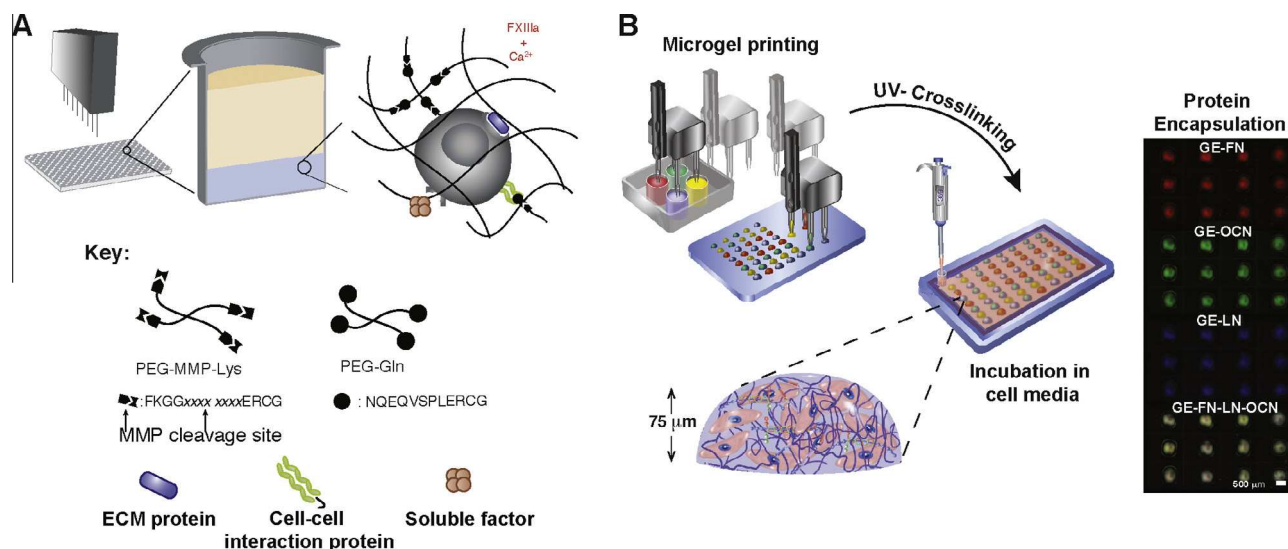


Fig. 5. Fabrication strategies of 3D microenvironments for stem cells. (A) Liquid handling robots dispense into 1536 microwell plates to generate a HTS platform of enzyme mediated cross-linked PEG gels that encompass tunable 3D microenvironments consisting of mESCs, degradable metalloproteinase (MMP) components, ECM proteins, cell-cell interaction proteins and soluble factors. (B) A gelatin methacrylamide solution containing hMSCs were contact printed and UV cross-linked to create 3D encapsulated hMSCs islands containing varying protein environments consisting of fibronectin (FN), osteocalcin (OCN) and laminin (LN). Adapted with permissions, (A) [76] 2014©, NPG (B) [77] 2014©, NPG.

In another approach, Khademhosseini and coworkers developed a combinatorial hydrogel microarray to investigate matrix effects on the osteogenesis of hMSCs [75]. Here, hMSCs were combined with liquid precursor solutions of a photoinitiator, methacrylated gelatin (GelMA), ECM proteins (fibronectin, laminin, and/or osteocalcin), and BMPs (BMP2 and/or BMP5) (Fig. 5B). The precursor solutions were contact printed onto functionalized glass slides and UV polymerized to form hydrogels. Alkaline phosphatase expression (a marker of osteogenesis) and alizarin red staining (an indicator of mineralization) were used as measures of the osteogenic potential of each material. In this study, hydrogels with all three ECM proteins (fibronectin, laminin, and osteocalcin) demonstrated improved osteogenesis as compared to controls and encapsulated ECM proteins had a more pronounced effect than soluble factors added to the medium.

5. Challenges accompanying 3D HTS

As 3D HTS techniques are adopted, there are several major challenges to consider. As cell biology moves to the third-dimension, new tools to understand and assess changes in material properties are required. As cells are increasingly cultured in degradable matrices, it is essential to develop imaging techniques to understand precisely how cells are altering the local environment over time. For example, micro-rheology has been used to monitor, in real-time, how a PEG based ECM mimic is remodeled during cell culture and migration [78]. In addition, cells push and pull on materials as they degrade and move through them and advances in quantifying these traction forces are needed. Toward this aim, imaging sensors have been designed to quantitate and understand cell-generated forces at the integrin level during adhesion and migration [79]. A DNA hairpin FRET-based probe was also developed to image in real-time cell generated forces at the cell–material interface [80].

Further, the biochemical signatures of the matrices are often altered during culture by deposition of proteins, from the cell culture medium and those expressed by the encapsulated cells. In order to quantify how cells interact with and sense their microenvironment, a more precise and comprehensive understanding of protein fouling and the evolution of the materials biochemical

landscape is needed. For example, improved mass spectrometry and elemental analysis methods are improving the detection of individual protein species on and within materials [81].

Simultaneously, improved techniques are necessary to elucidate cell phenotype in 3D. Advances in reporter cell lines have aided [74], but there is still a need to develop easy readouts of cell function that can be used to correlate material properties to cell behavior that are amenable to HTS. This includes new probes for monitoring intracellular signaling, cytokine secretion, differentiation, surface receptor presentation and binding, as well as transcriptional landscapes. Many of the standard molecular biology techniques become significantly more complicated in 3D and during HTS. In order for 3D HTS to be adopted beyond very preliminary screening, the field will need to adapt molecular biology approaches to these more complex settings.

Finally, the ECM is not a static material and many cues that are important in development and disease progression alter in time [61]. Dynamic biomaterials that enable user-defined control over material properties to study how a stiffening matrix influences fibrosis or growth factor secretion regulates neurogenesis are needed [82]. Ultimately, matrices that enable reversible biophysical and biochemical properties will be needed to capture the full dynamics of the native ECM as seen in development and disease. This has been captured using photoresponsive azobenzene crosslinks [83] and sequential photoaddition and photocleavage reactions [84]. All of these techniques become increasingly difficult to incorporate into HTS approaches, but the chemical advances in the biomaterials field are bringing these concepts into reality.

6. Summary

The studies discussed in this review have identified peptide ligands, chemical moieties, elastic moduli and topographical patterns that influence the behavior of stem cells. It is envisioned that future HTS platforms will advance further to incorporate dynamic biomaterials for investigation of spatiotemporal effects. As we look ahead, combinations of 2D and 3D HTS approaches will be indispensable in providing critical information as to how the environment informs stem cell function and fate. HTS studies have

helped to gain retrospective mechanistic insights and it is hoped that further development of high content analytical tools will allow elucidation of the role of material properties identified during screening to help build roadmaps that will instruct materials development for regenerative medicine applications such as *ex vivo* tissue growth, stem cell maintenance and controlled expansion.

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